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Identification Scheme of Table 1 Biomarkers

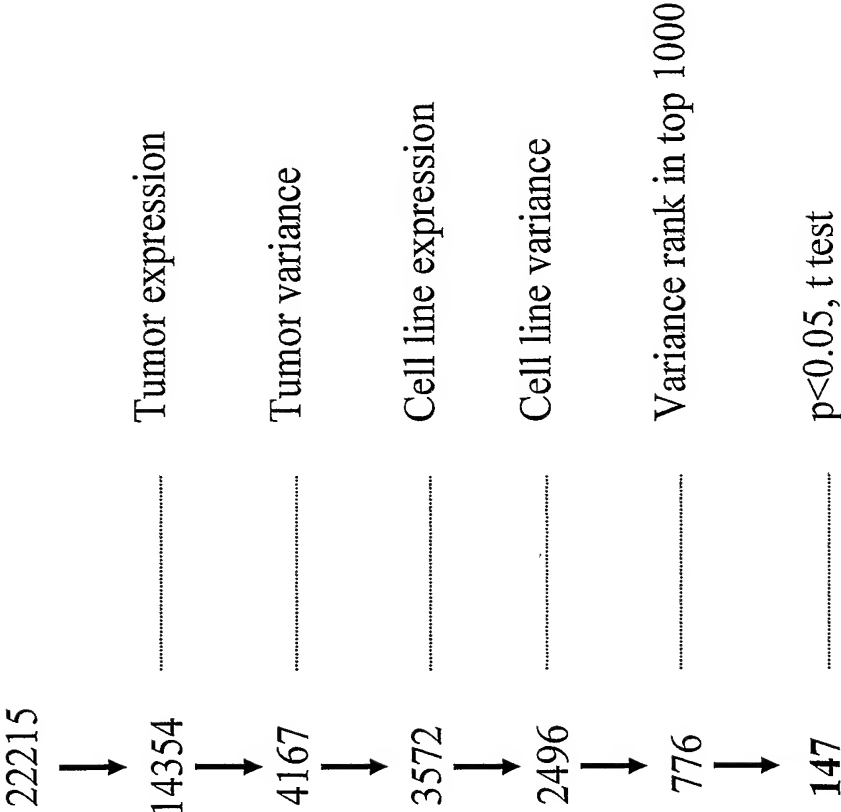


FIG. 1

Identification Scheme of Table 2 Biomarkers

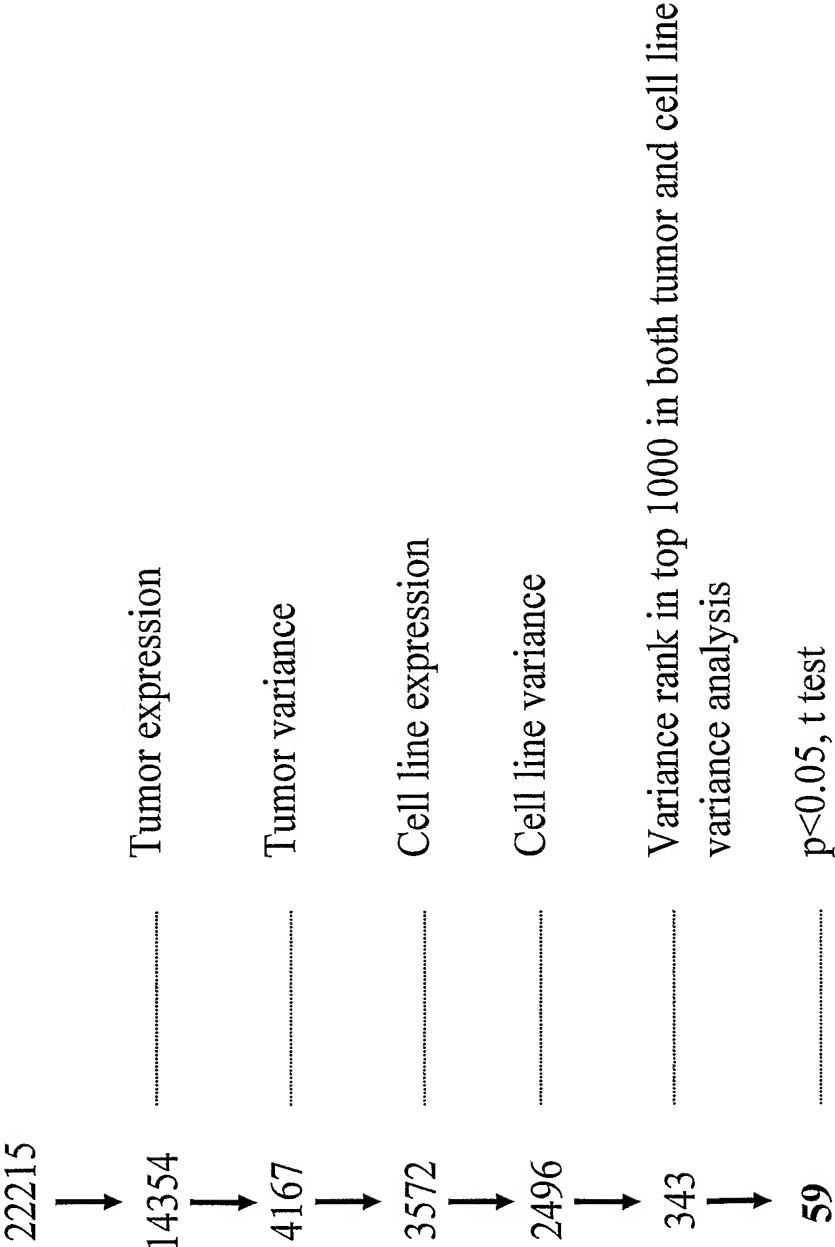


FIG. 2

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EGFR Expression in NSCLC Cell Lines

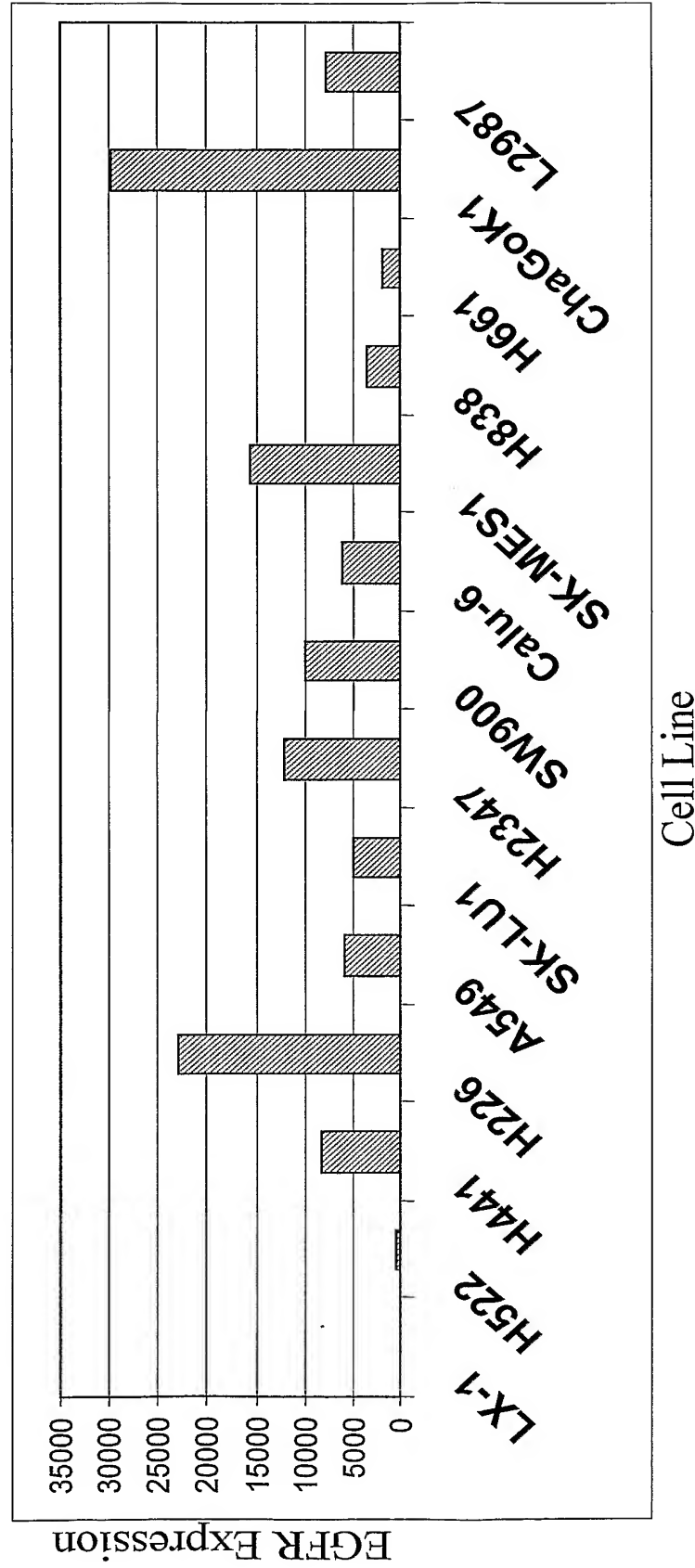


FIG. 3

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Variance Analysis of Expression Profiles

29 Adenocarcinoma Tumors

- 22,215 probe set Affymetrix GeneChip® HG-U133A array profiles of 29 lung adenocarcinoma tumors
- 14,354 probe sets present ($p \leq 0.04$) in at least 2 samples
- Calculate variance and rank probe sets

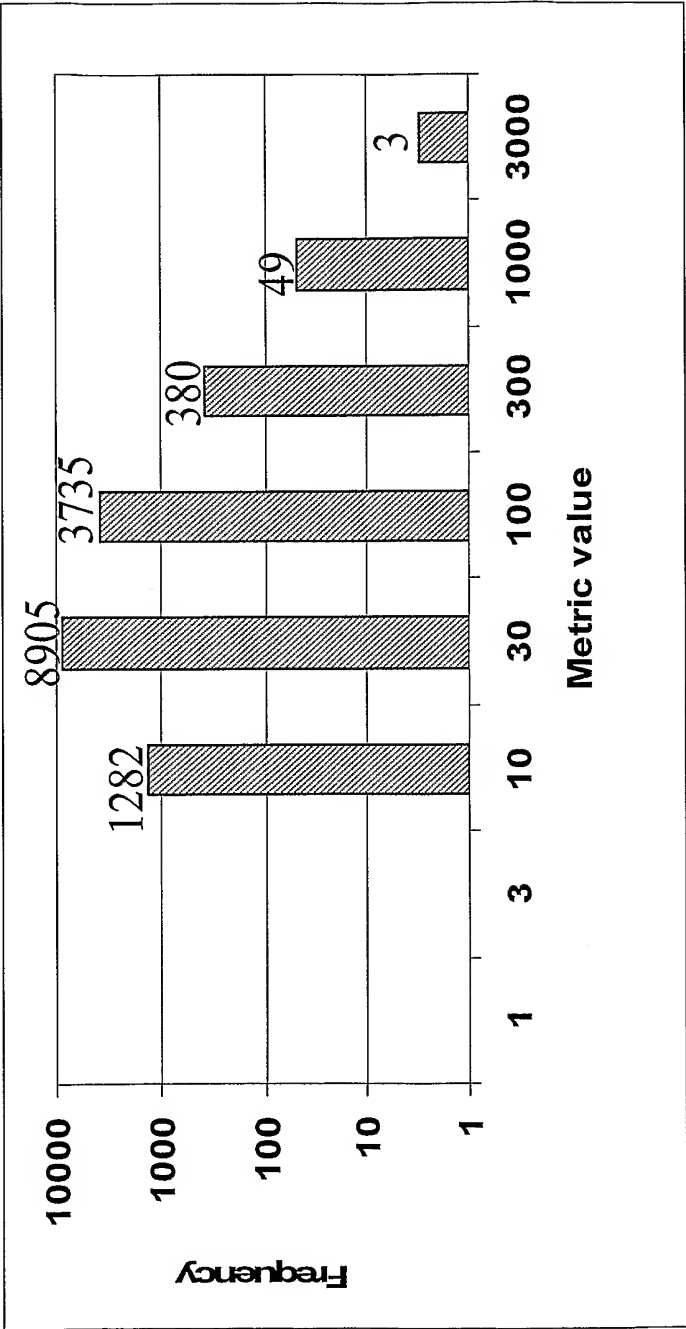
14 NSCLC Cell Lines

- 22,215 probe set Affymetrix GeneChip® HG-U133A array profiles of 14 NSCLC cell lines
- 13,909 probe sets present ($p \leq 0.04$) in at least 2 cell lines
- Calculate variance and rank probe sets

FIG. 4

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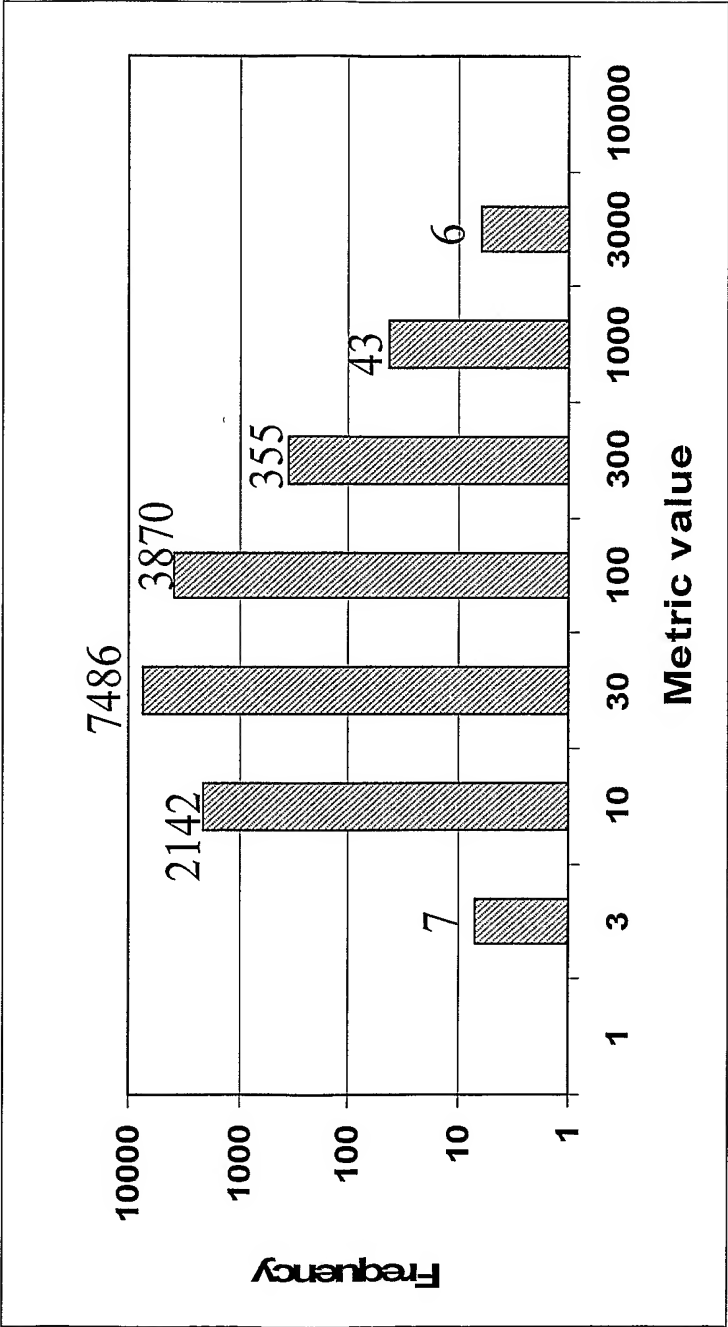
Variance Metric Distribution of Probe sets



Adenocarcinoma Tumors - 4167 probe sets of WSpread(90-10) ≥ 30

FIG. 5

Variance Metric Distribution of Probe sets

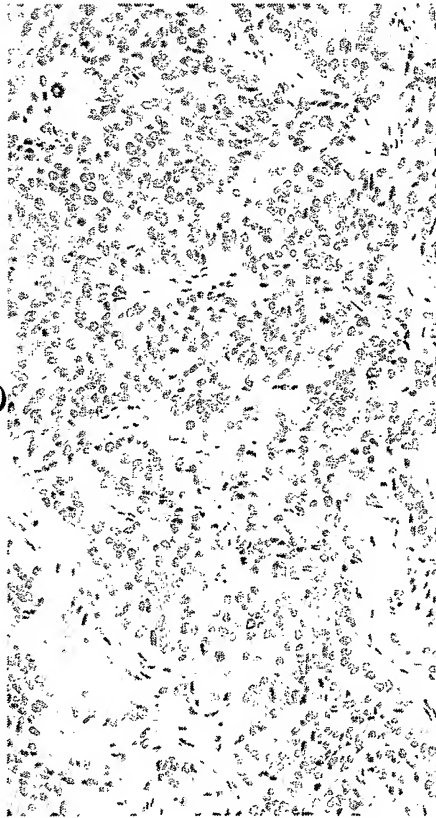


Cell Lines - 4274 probe sets of WSpread(90-10) ≥ 30

FIG. 6

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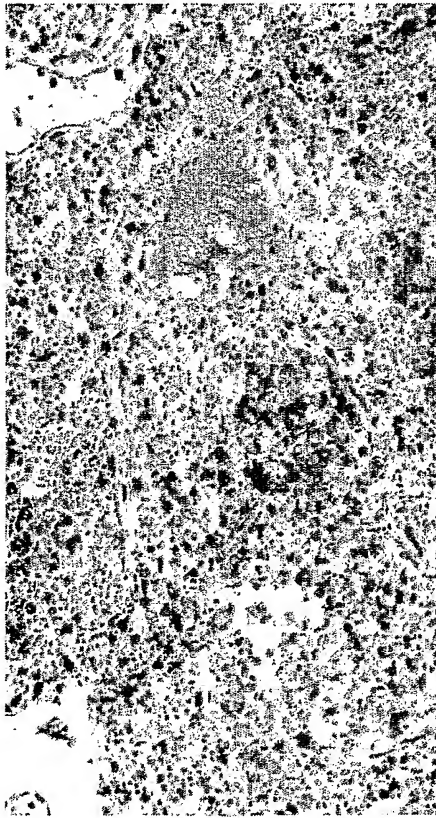
0 Negative



0.5 Low



1+ Moderate



2+ High



FIG. 7